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In the claims

Please amend claims 5 and 6 to read as follows:

5. (Amended) A method for determining the phenotype of a test cell from a given tissue, comprising detecting the presence or absence of differential expression, relative to a normal cell of the given tissue type, of at least 5 different genes shown in Table 1,

wherein the presence of differential expression indicates that said test cell has an IBD or pre-IBD phenotype.

6. (Amended) The method of claim 5, wherein said differential expression is upregulation or downregulation by at least a factor of two.

Please add the following new claims:

- --19. (New) The method of claim 5, wherein said test cell is an intestinal cell.
- 20. (New) The method of claim 5, comprising detecting the presence or absence of differential expression of at least 10 different genes shown in Table 1.
- 21. (New) The method of claim 5, comprising detecting the presence or absence of differential expression of at least 25 different genes shown in Table 1.

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- 22. (New) The method of claim 5, comprising detecting the presence or absence of differential expression of at least 50 different genes shown in Table 1.
- 23. (New) The method of claim 5, comprising detecting the presence or absence of differential expression of at least 75 different genes shown in Table 1.
- 24. (New) The method of claim 5, wherein said genes belong to distinct functional classes.
- 25. (New) The method of claim 5, wherein said detecting comprises in situ hybridization.
- 26. (New) The method of claim 5, wherein said detecting comprises hybridization to nucleic acid probes immobilized on a solid support.
- 27. (New) The method of claim 26, wherein said nucleic acid probes are immobilized in a two-dimensional array.
- 28. (New) A method for determining the phenotype of a test cell from a given tissue, comprising detecting the presence or absence of differential expression, relative to a normal cell of the given tissue type, of at least 5 different genes shown in Table 1, said genes belonging to distinct functional classes,

wherein the presence of differential expression indicates that said test cell has an IBD or pre-IBD phenotype.

